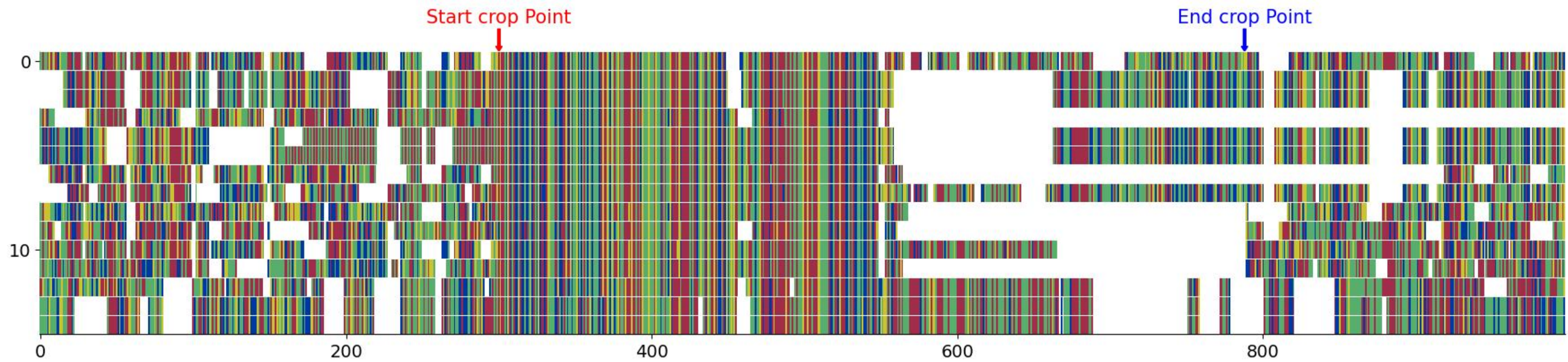
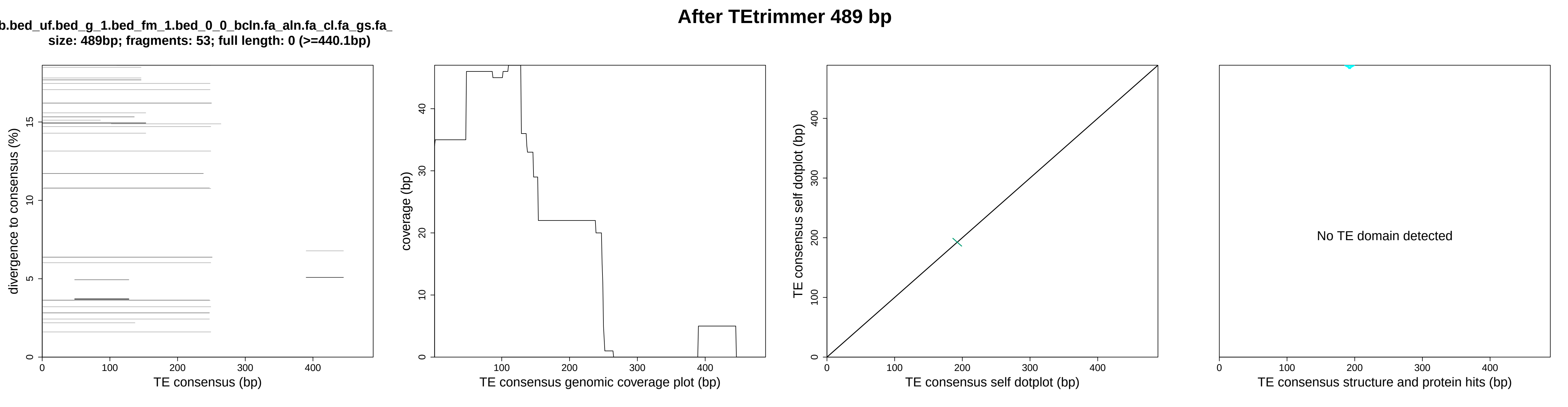


MSA length = 489

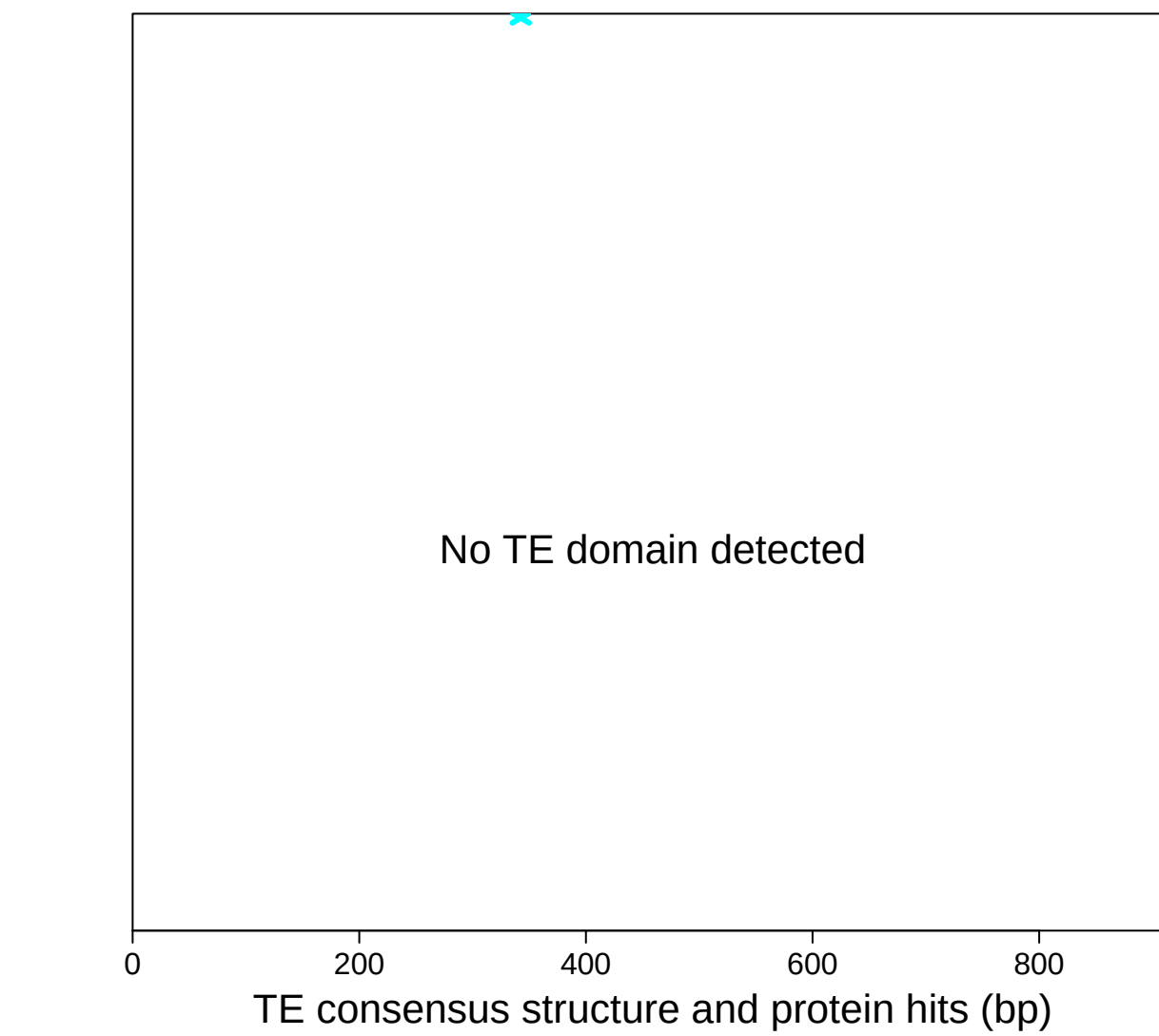
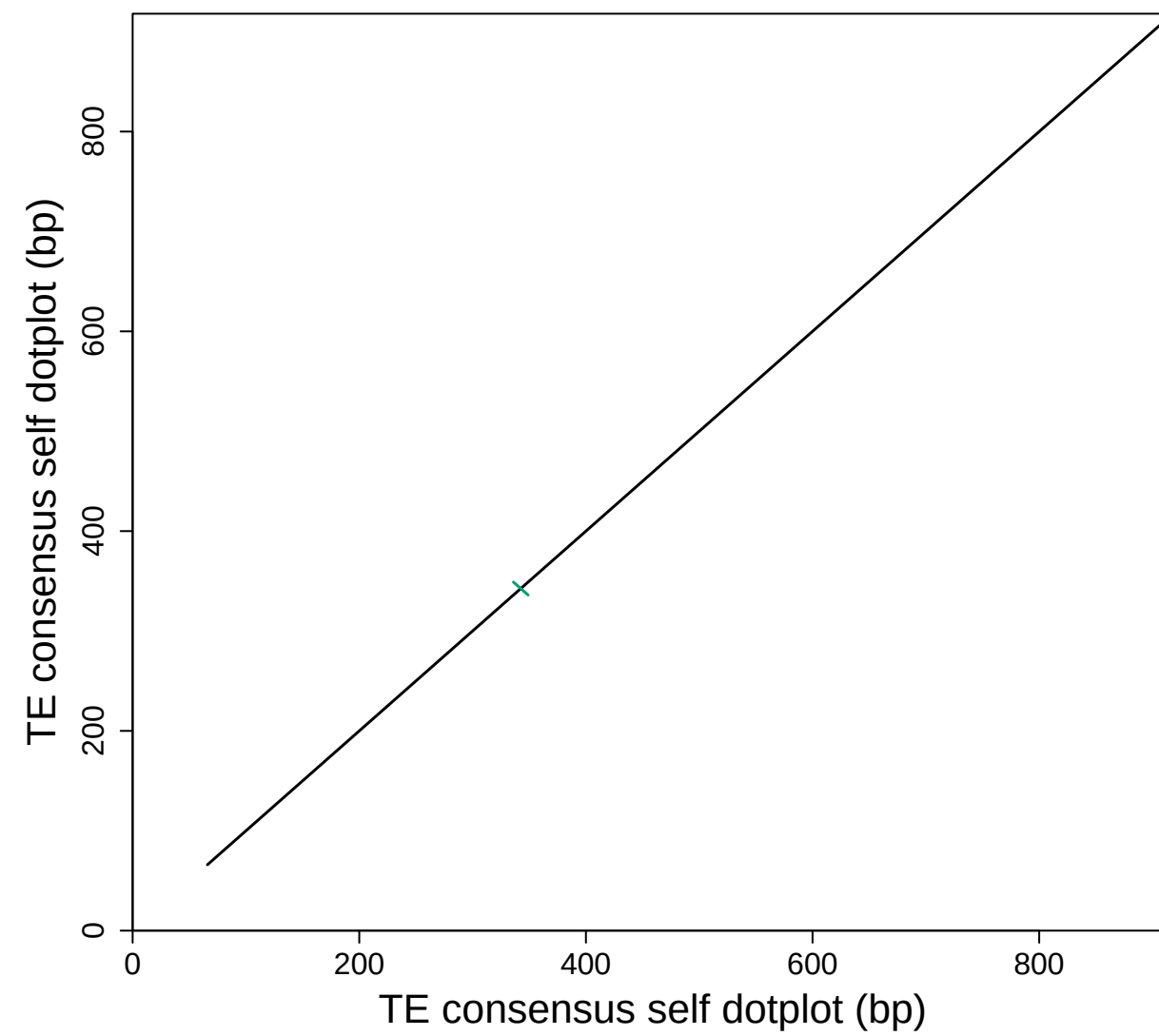
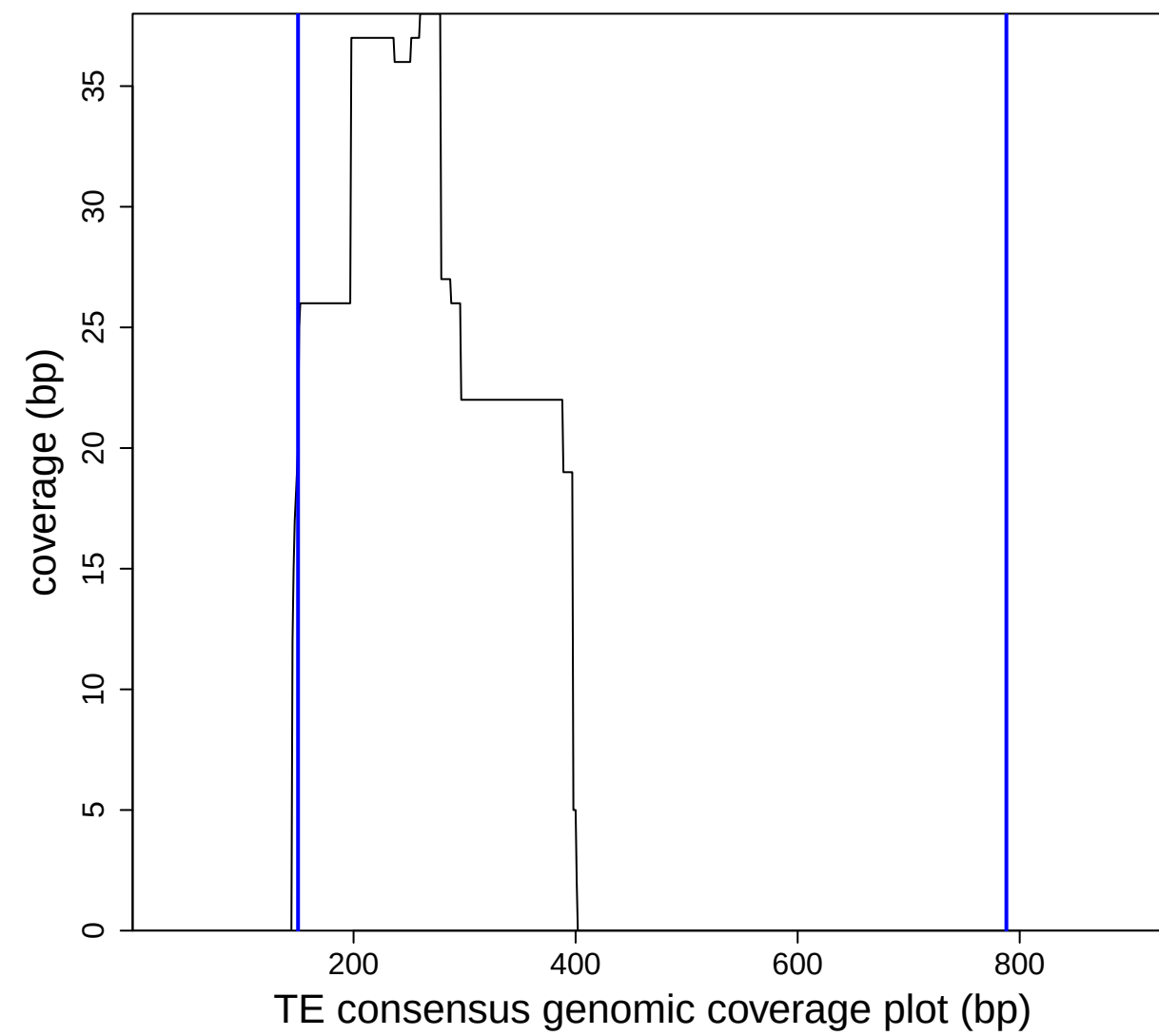
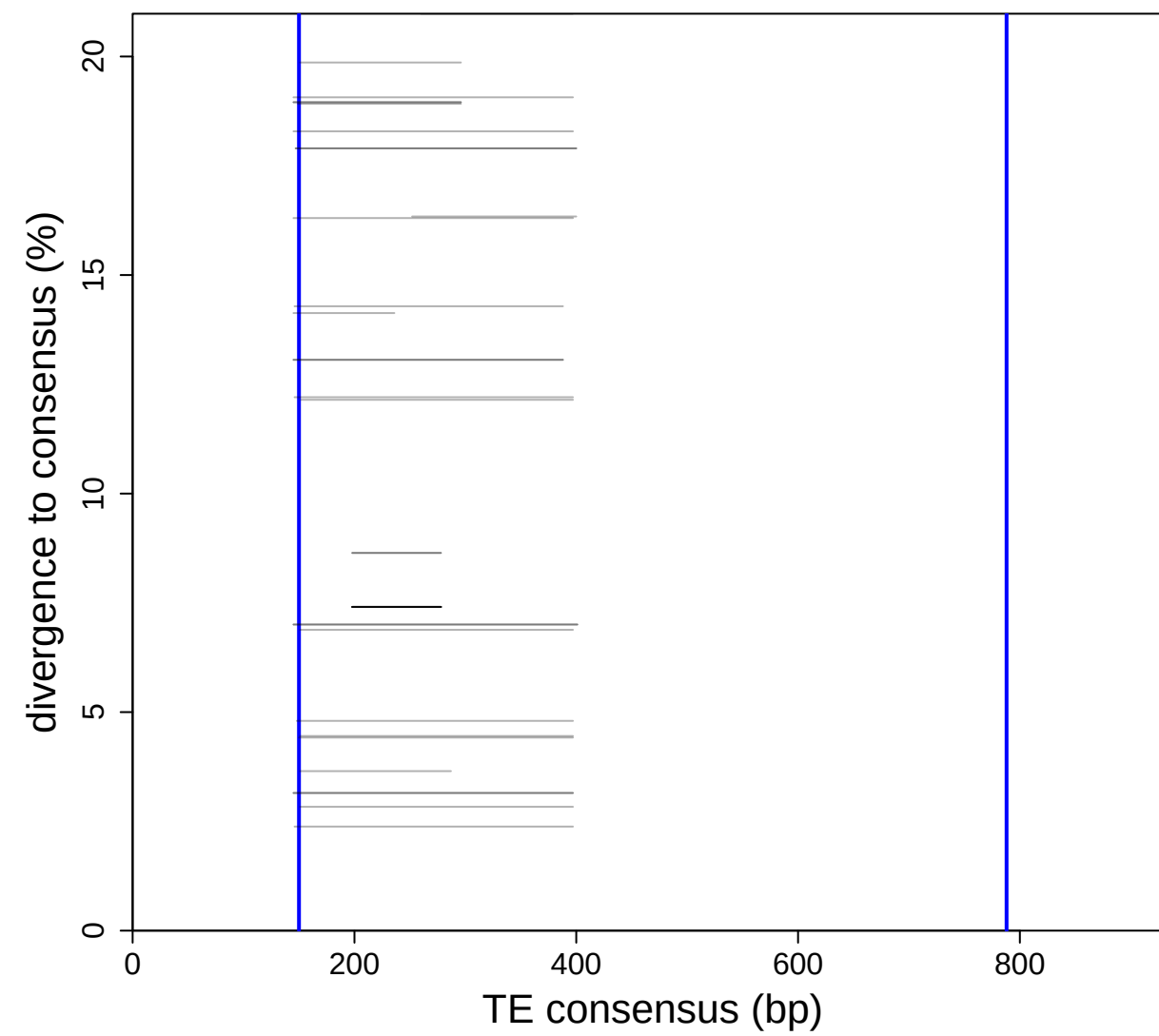
[illegible]



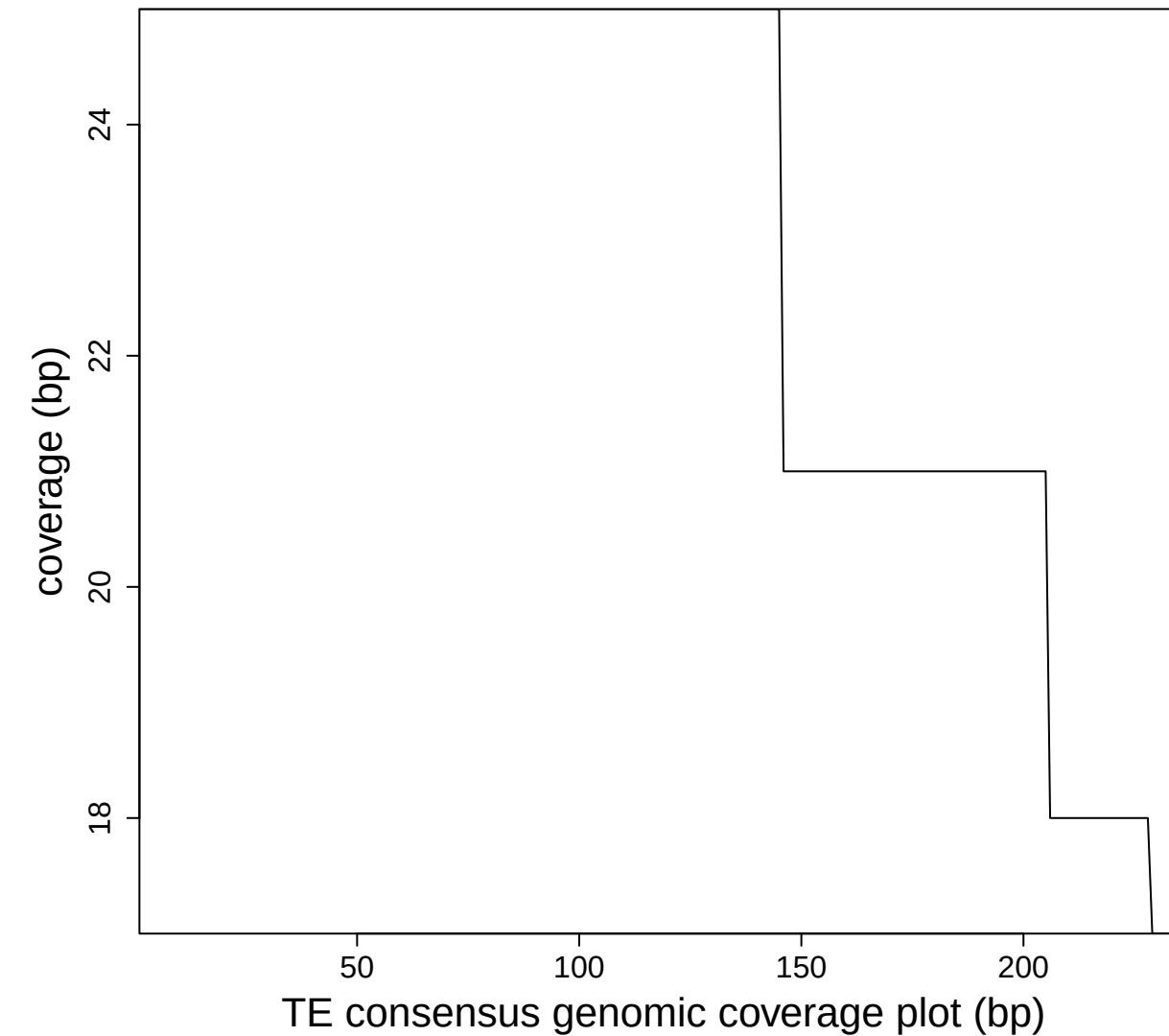
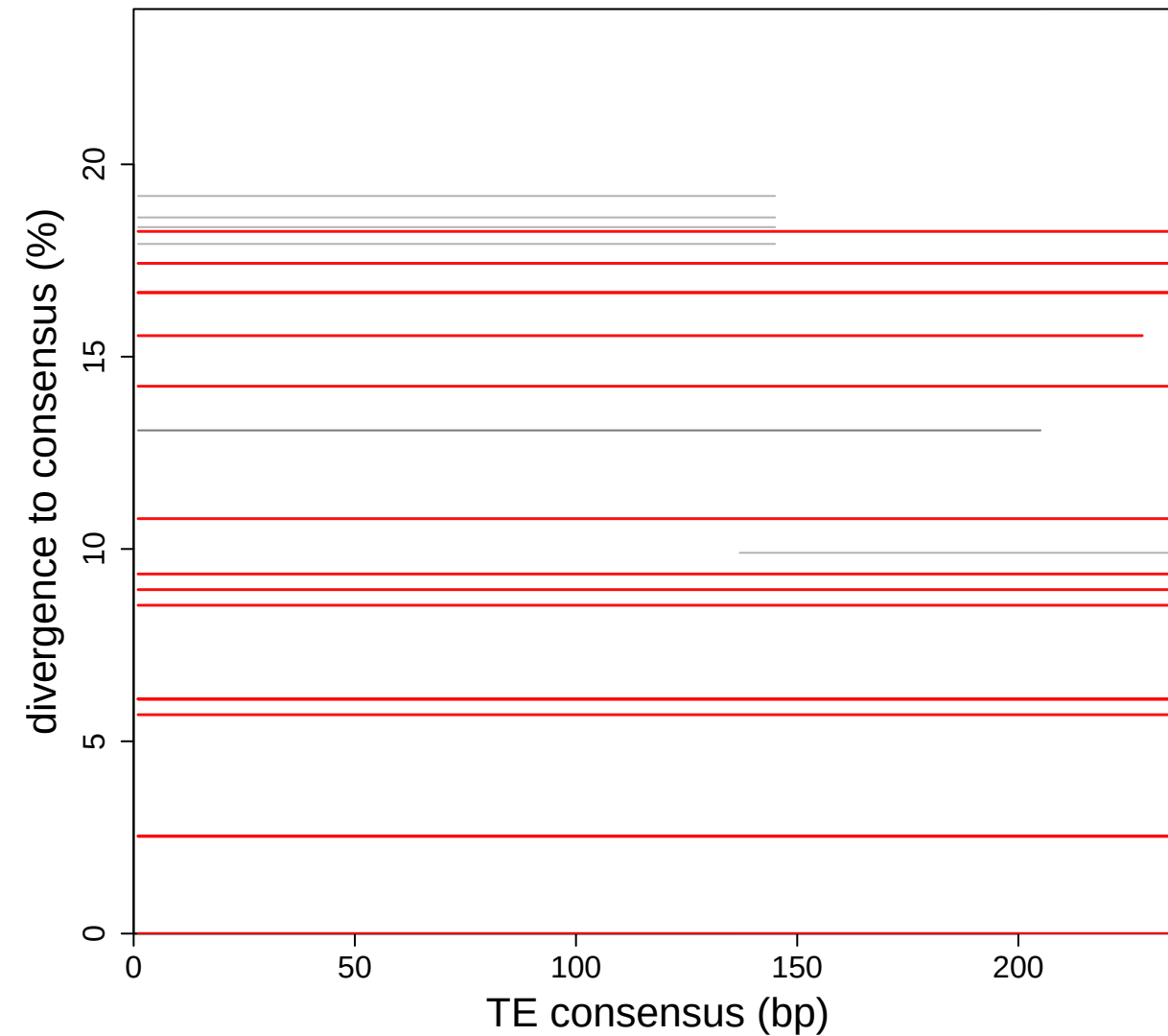


fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa_
size: 938bp; fragments: 39; full length: 0 (>=844.2bp)

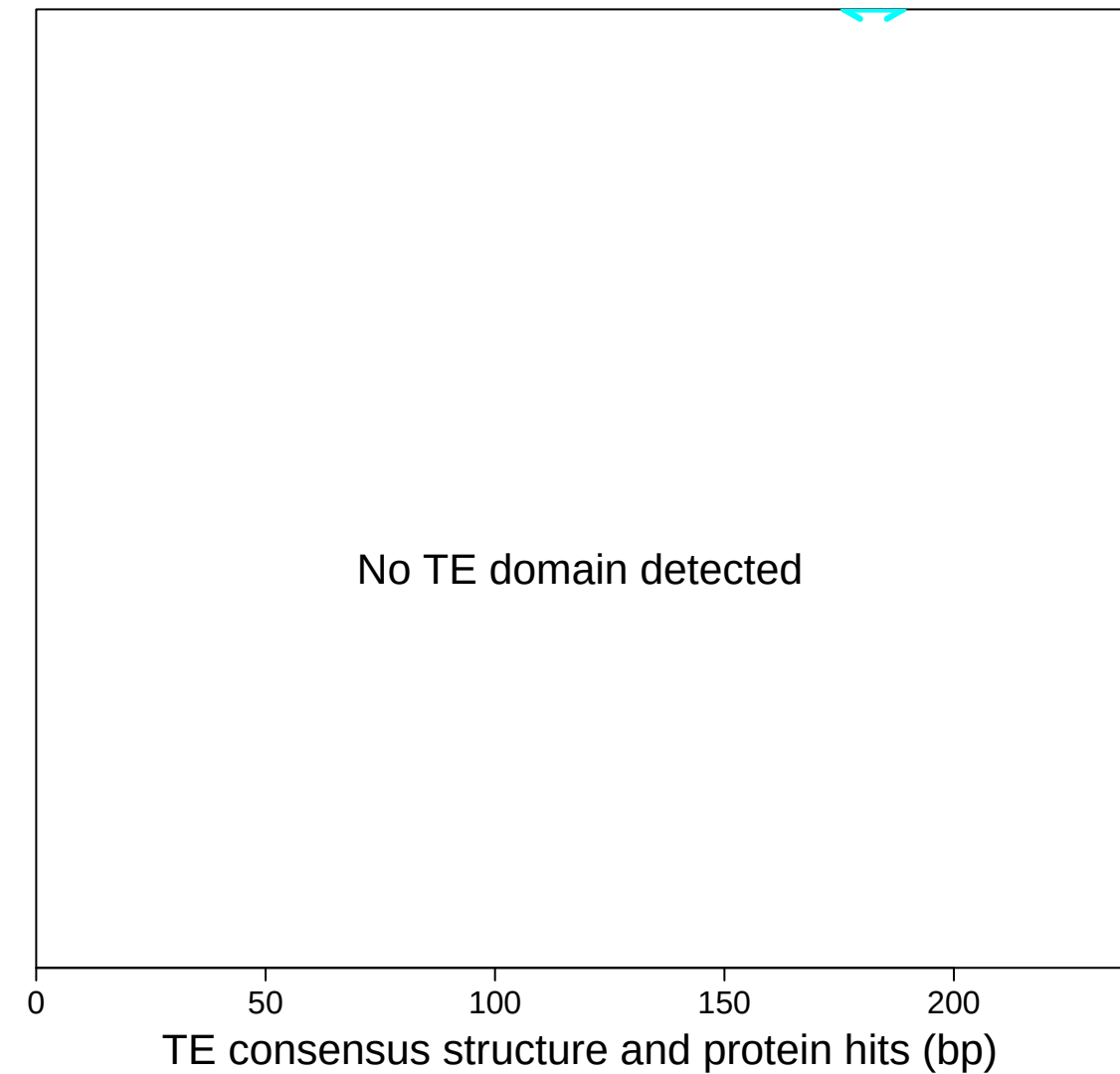
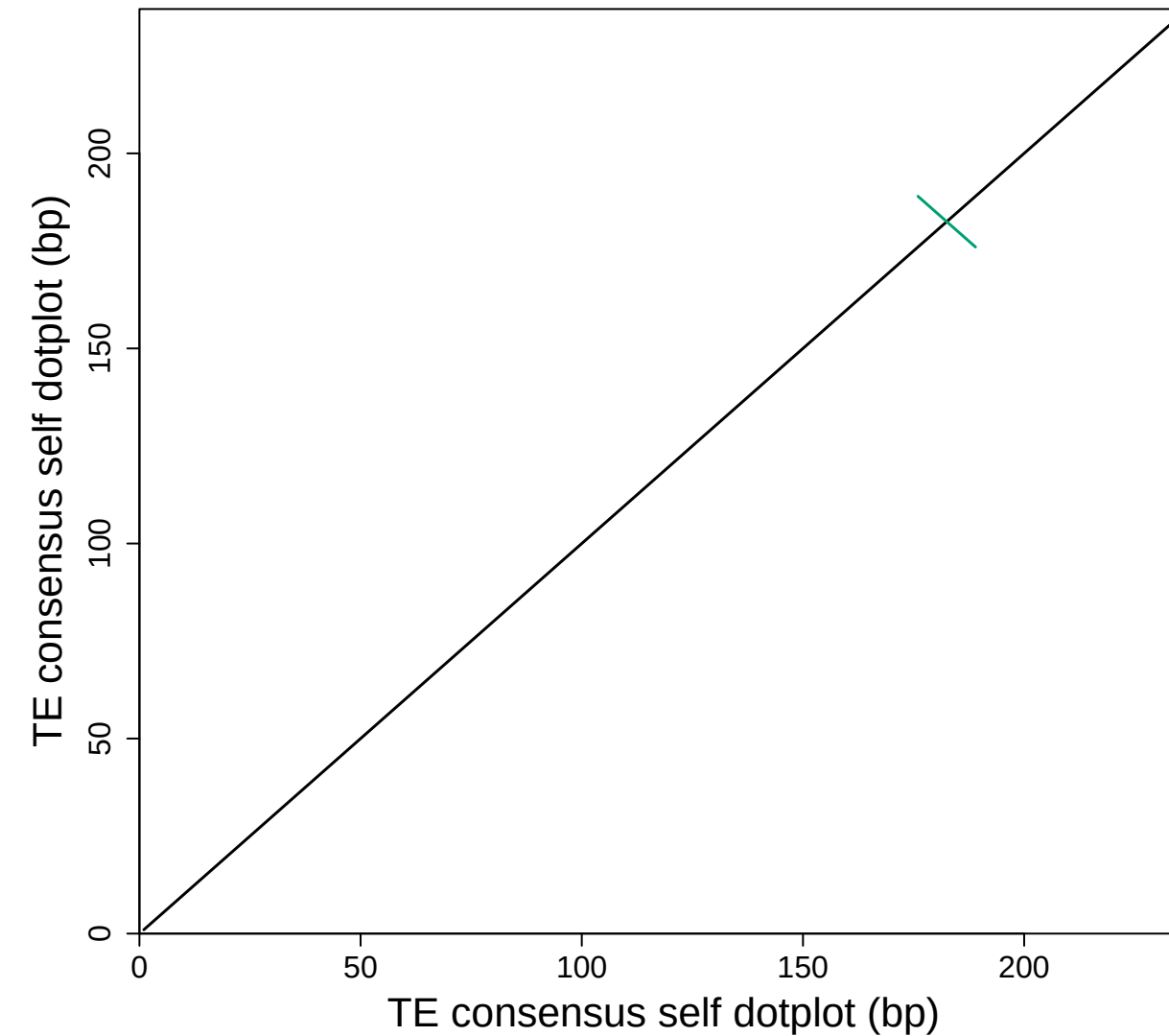
After TEtrimmer Extended plot Blue lines are boundaries



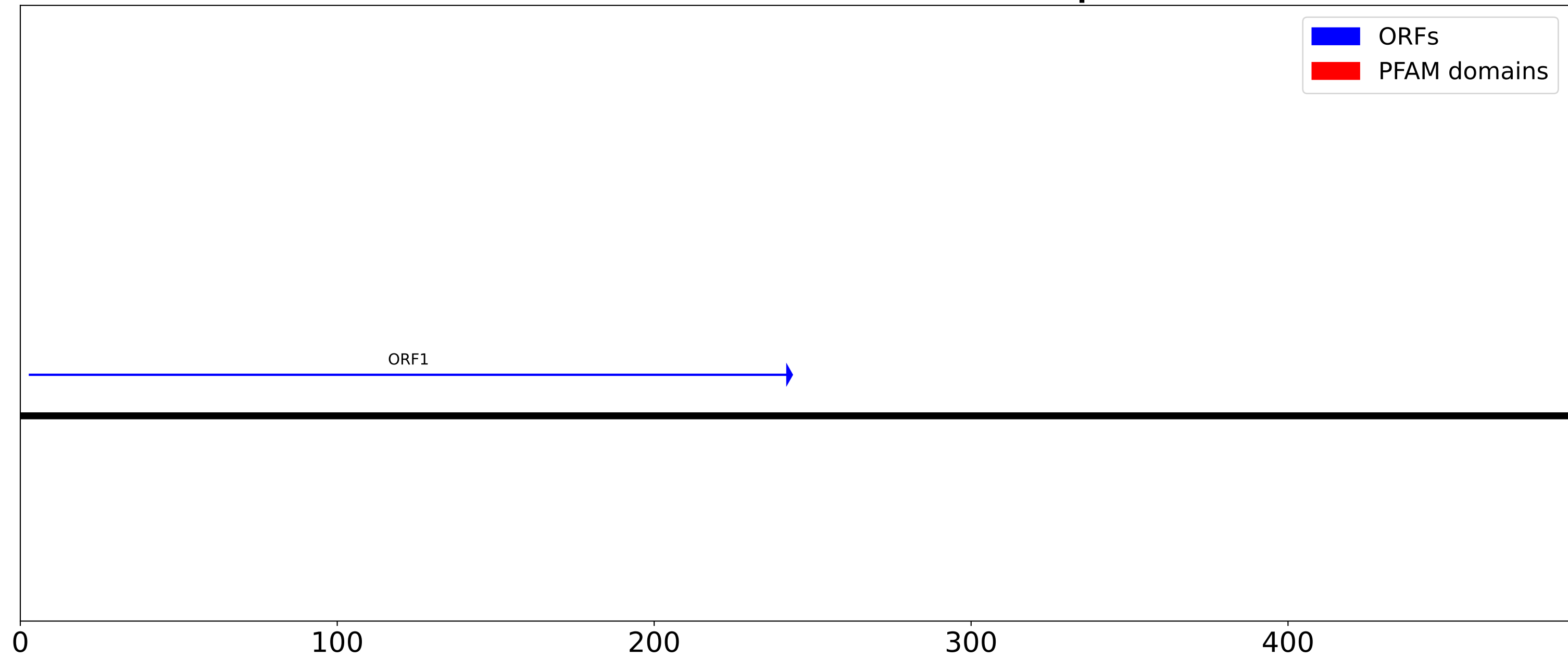
TE: ltr_1_family_12
size: 237bp; fragments: 26; full length: 17 (≥ 213.3 bp)



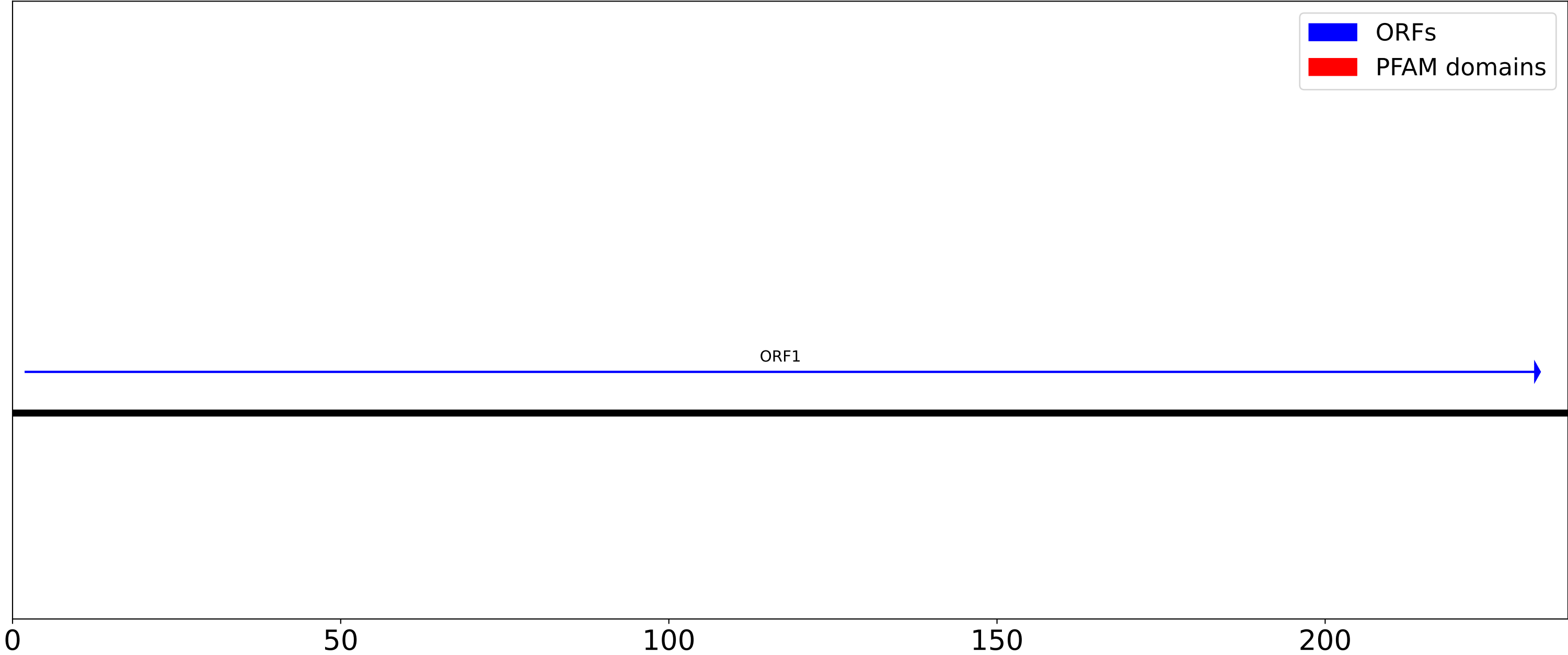
Before TEtrimmer 237 bp



After TEtrimmer ORF and PFAM domain plot



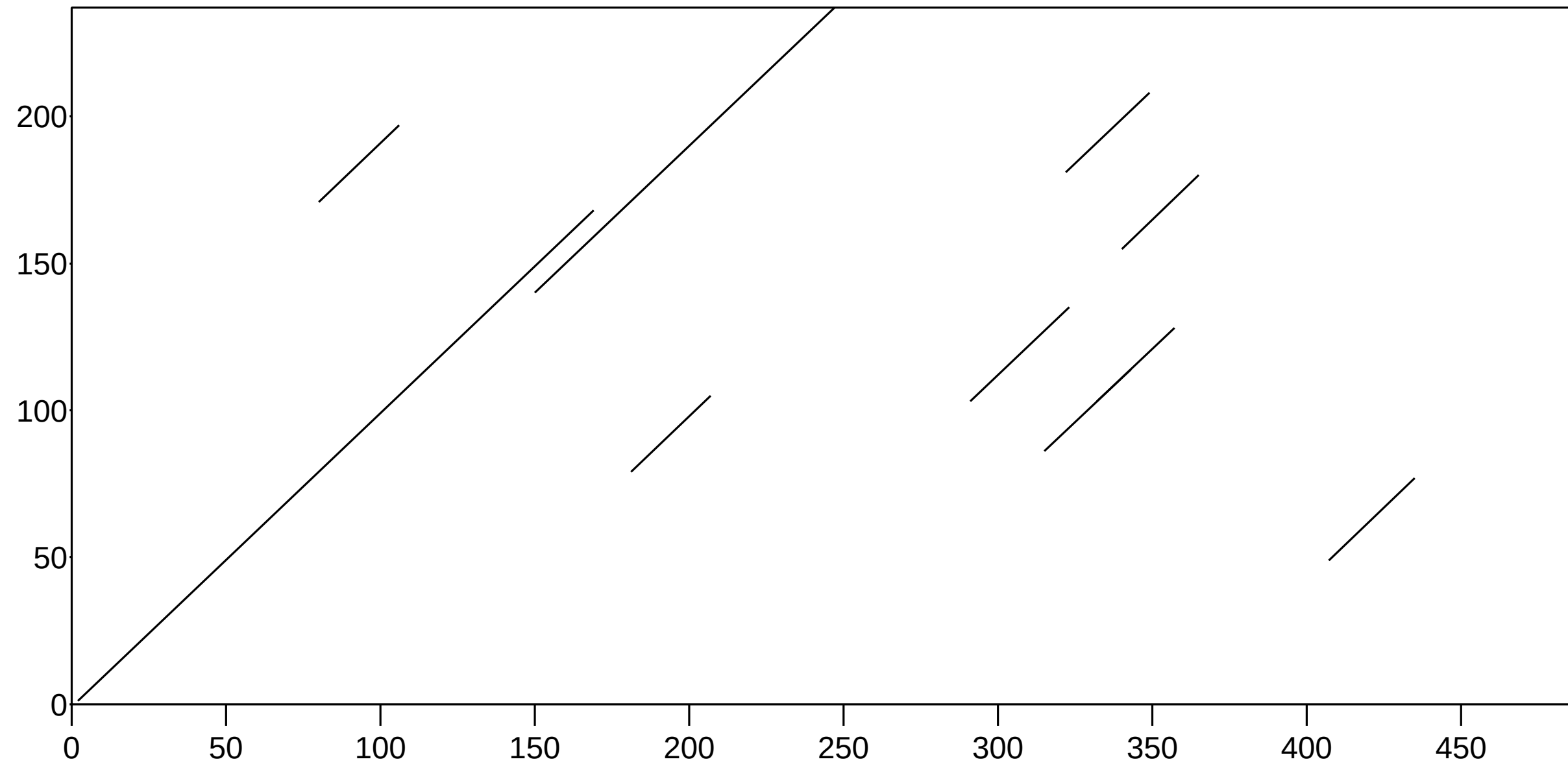
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)